SNCBI	250		ے Nucleo	tide		
PubMed No	ucleotide Protein	Genome S	tructure PMC	Taxonomy	OMIM	Books
Search Nucleotide	▼ for				Go Clear	
	Limits	Preview/Index	History	Clipboard	De	tails
Display default	▼ Show:20	▼ Send to File				
1: BQ072384. AGENCOURT_6838696[gi:19901430]						Links
IDENTIFIERS						
dbEST Id: EST name: GenBank Acc: GenBank gi:	11922326 AGENCOURT_683 BQ072384 19901430	8696				

CLONE INFO

Clone Id:

IMAGE:5761537 (5')

Plate: LLAM12810 Row: f Column: 02

DNA type: cDNA

PRIMERS

PolyA Tail: Unknown

SEQUENCE

TTTTTTTCCCCCCCCCCAAAGAGGGGGTTTTTTTTT

Quality:

High quality sequence stops at base: 232

Entry Created: Last Updated:

Mar 28 2002 Apr 2 2002

COMMENTS

Tissue Procurement: Life Technologies, Inc.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Agencourt Bioscience Corporation

Clone distribution: MGC clone distribution information can

be found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llnl.gov

LIBRARY

Lib Name: NIH_MGC_122
Organism: Homo sapiens

Organ: pooled lung and spleen

Lab host: DH10B

Vector: pCMV-SPORT6

R. Site 1: NotI

R. Site 2: Description: EcoRV (destroyed)

RNA source anonymous pool of 24 week female lung, 16 week female spleen, and 20-22 week male spleens. Library is

female spleen, and 20-22 week male spleens. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.4 kb, insert size range 1-3 kb. Library is normalized and enriched for

full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 026. Note:

this is a NIH_MGC Library.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Institutes of Health, Mammalian Gene Collection

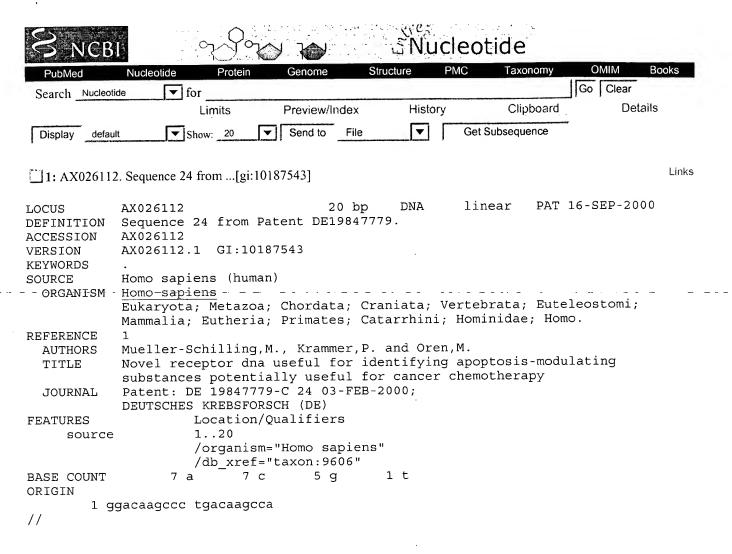
(MGC)

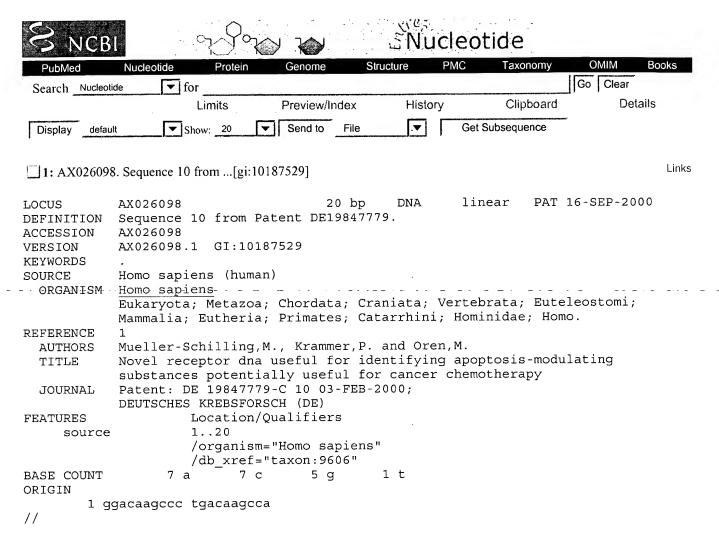
Authors: NIH-MGC http://mgc.nci.nih.gov/

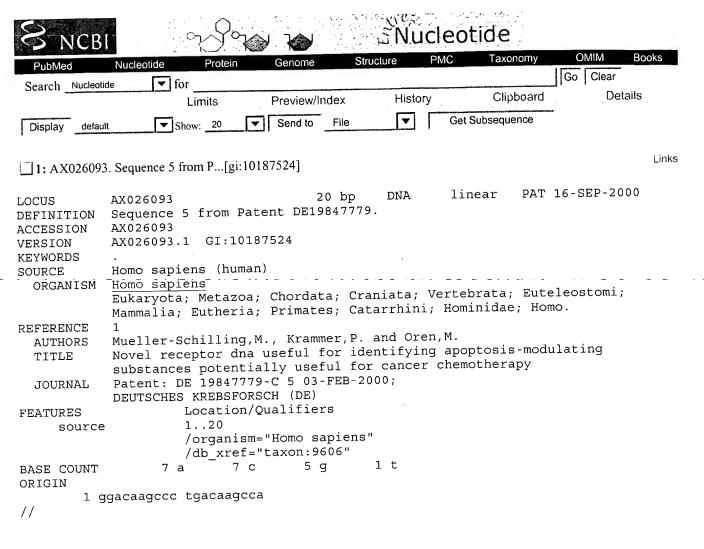
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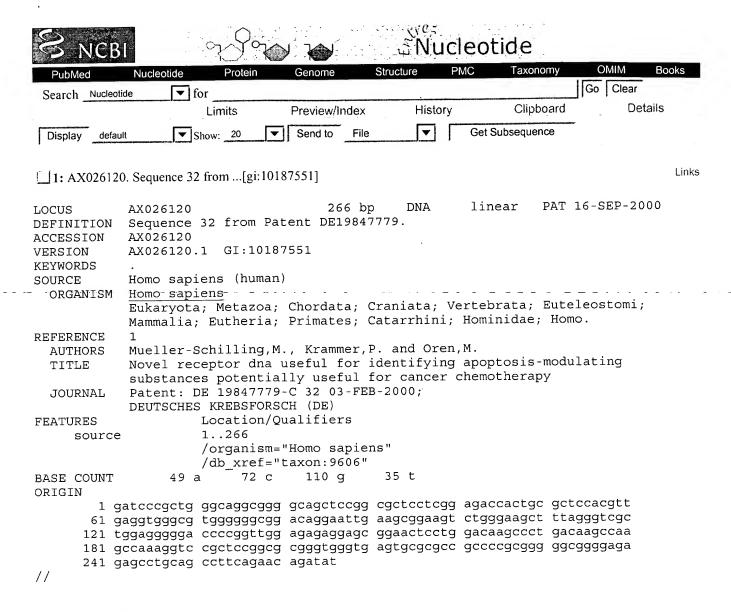
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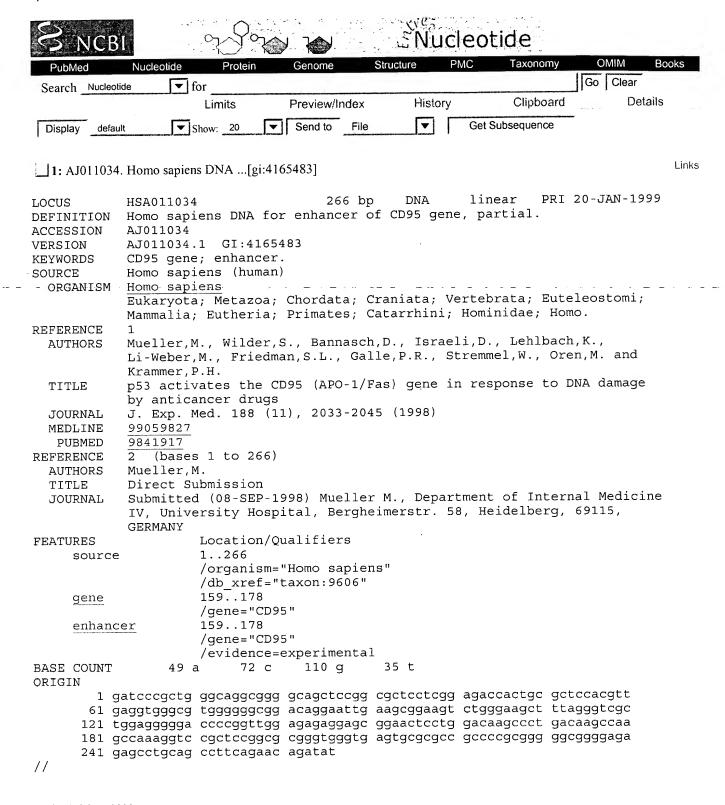
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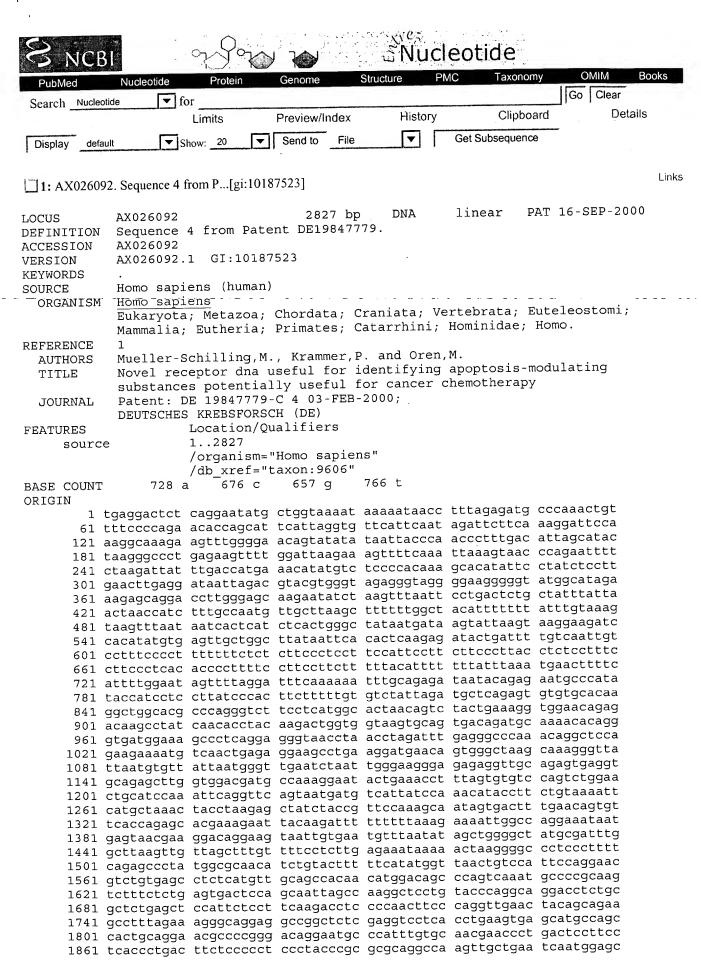




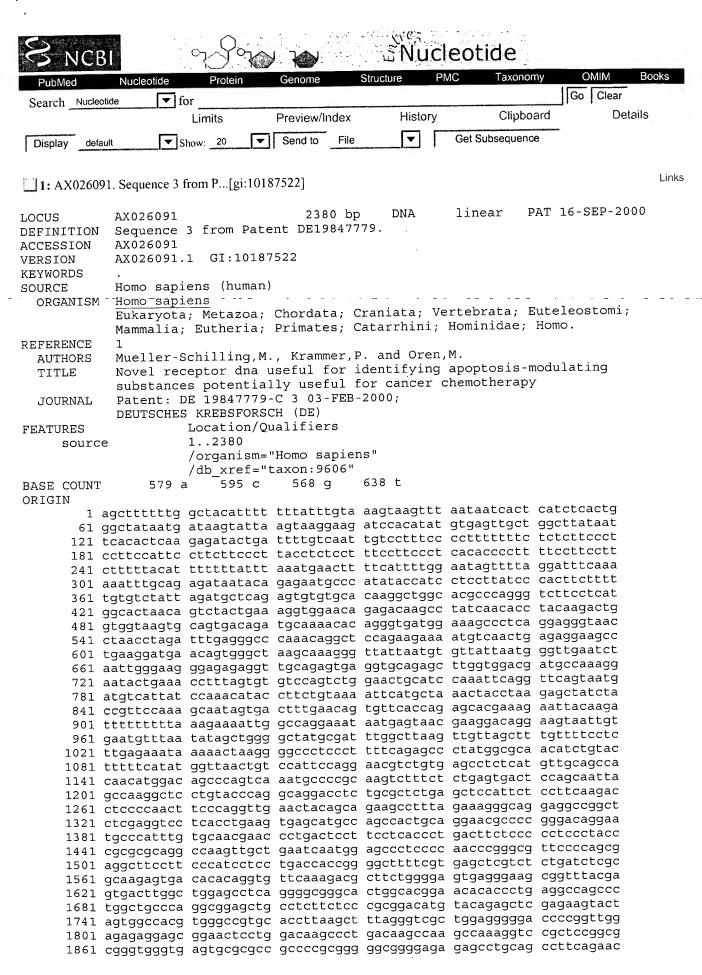






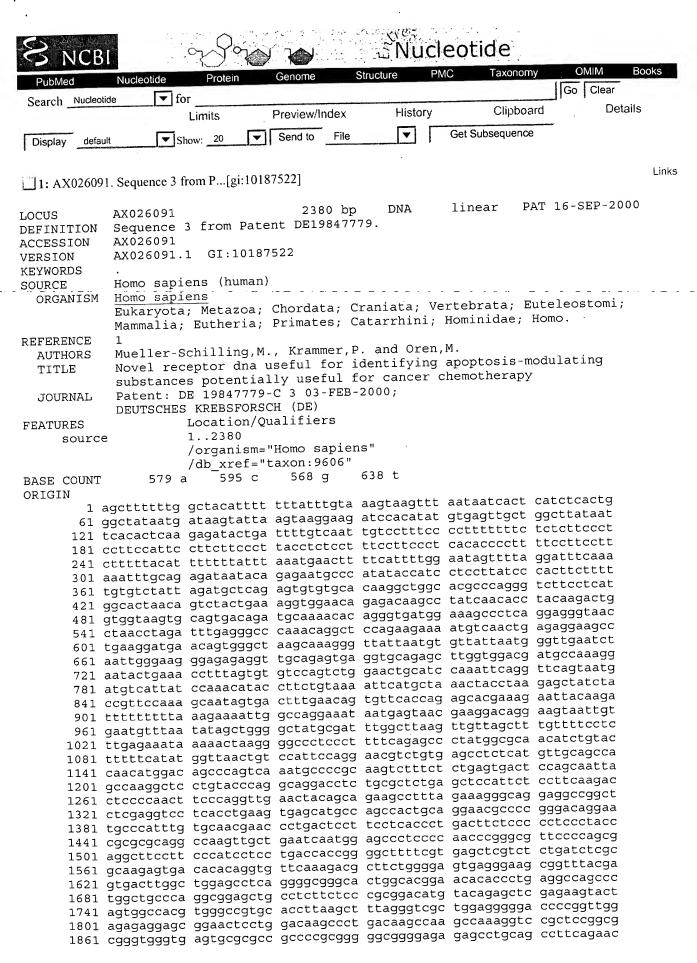


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    2101 gcacggaaca caccetgagg ccagceetgg etgeccagge ggagetgeet etteteeege
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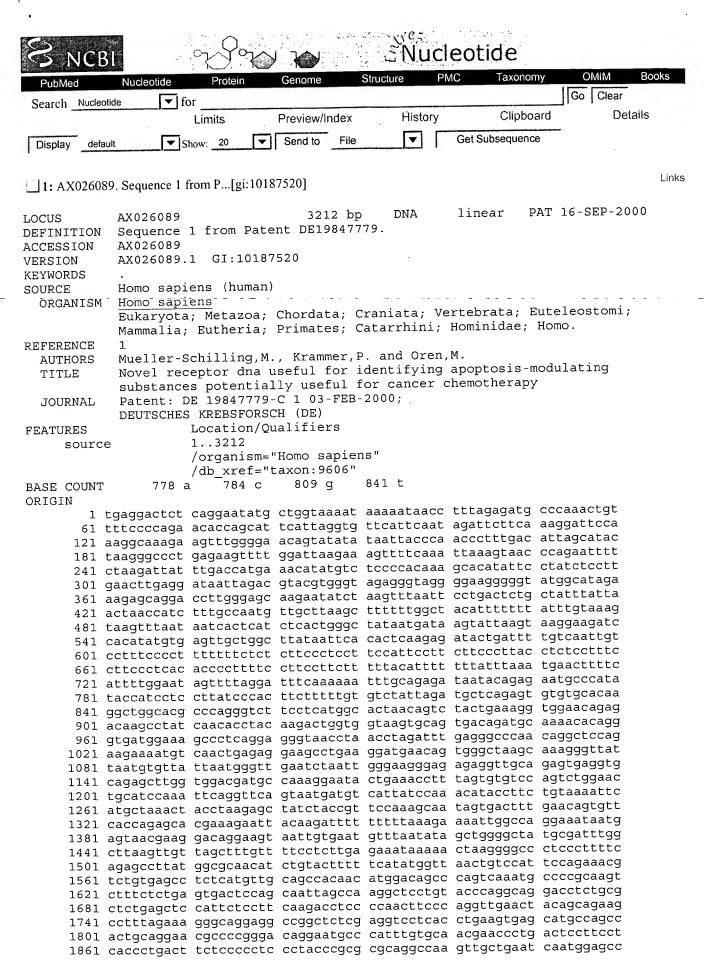


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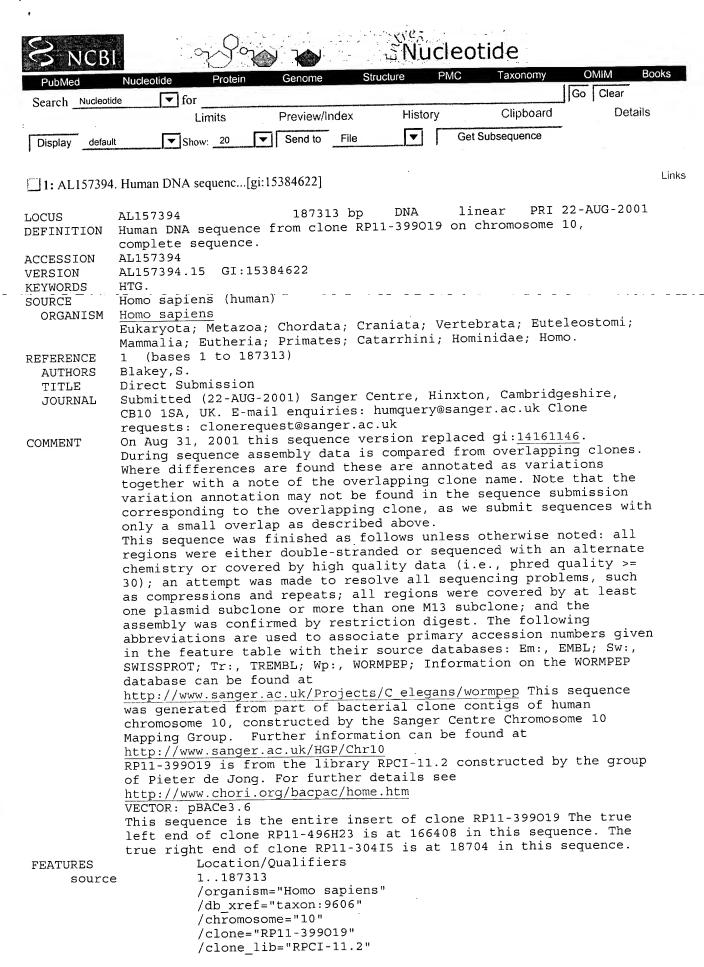


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    2041 tggggagtga gggaagcggt ttacgagtga cttggctgga gcctcagggg cgggcactgg
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    DNA (human gene TNFRSF6) (9CI) (CA INDEX NAME)
CN
OTHER NAMES:
                                                    703
    310: PN: WO03008583 TABLE: 54 claimed DNA
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FS
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CI
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SR
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     STN Files: CA, CAPLUS, TOXCENTER
LC
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L1
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 SR
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 LC
 **RELATED SEQUENCES AVAILABLE WITH SEQLINK**
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 *** USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE ***
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LC
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
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**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
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L1
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RN
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     complementary (9CI) (CA INDEX NAME)
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DNA, d(T-G-G-C-T-T-G-T-C-A-G-G-G-C-T-T-G-T-C-C), double-stranded

OTHER CA INDEX NAMES:

Unspecified

OTHER NAMES:

MAN

CA

FS

MF CI

SR

LC

complementary (9CI)

NUCLEIC ACID SEQUENCE

STN Files: CA, CAPLUS

2: PN: DE19847779 FIGURE: 4 claimed DNA

ANSWER 14 OF 14 REGISTRY COPYRIGHT 2003 ACS 220304-11-0 REGISTRY L1

RN

DNA (human gene CD95 p53-binding site-containing fragment) (9CI) (CA CN INDEX NAME)

OTHER NAMES:

1: PN: DE19847779 FIGURE: 4 claimed DNA CN

DNA (human Fas gene promoter region tumor antigen p53-responsive CNelement-containing fragment)

NUCLEIC ACID SEQUENCE FS

Unspecified MF

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER

RELATED SEQUENCES AVAILABLE WITH SEQLINK

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